

SCORE Search Results Details for Application 10687035 and Search Result 20080310_104727_us-10-687-035-33.rai.

Score Home	Retrieve Application	SCORE System	SCORE	Comments /
Page	List	Overview	FAQ	Suggestions

This page gives you Search Results detail for the Application 10687035 and Search Result 20080310_104727_us-10-687-035-33.rai.

[Go Back to previous page](#)

GenCore version 6.2.1

Copyright (c) 1993 - 2008 Bioceleration Ltd.

OM protein - protein search, using sw model

Run on: March 10, 2008, 14:04:05 ; Search time 36 Seconds
(without alignments)
557.713 Million cell updates/sec

Title: US-10-687-035-33
Perfect score: 656
Sequence: 1 MDFQVQIFSFLISASVIMS.....YCQQWSSNPFTFGSGTKLEI 127

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1048630 seqs, 157249835 residues

Total number of hits satisfying chosen parameters: 1048630

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_AA:*
1: /ABSS/Data/CRF/ptodata/2/iaa/5_COMB.pep:*
2: /ABSS/Data/CRF/ptodata/2/iaa/6_COMB.pep:*
3: /ABSS/Data/CRF/ptodata/2/iaa/7_COMB.pep:*
4: /ABSS/Data/CRF/ptodata/2/iaa/H_COMB.pep:*
5: /ABSS/Data/CRF/ptodata/2/iaa/PCTUS_COMB.pep:*
6: /ABSS/Data/CRF/ptodata/2/iaa/RE_COMB.pep:*
7: /ABSS/Data/CRF/ptodata/2/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

%
Result Query

No.	Score	Match	Length	DB	ID	Description
1	572	87.2	128	1	US-08-476-275-4	Sequence 4, Appli
2	572	87.2	128	2	US-08-475-815B-7	Sequence 7, Appli
3	572	87.2	128	2	US-08-475-813-4	Sequence 4, Appli
4	572	87.2	128	3	US-10-411-037-60	Sequence 60, Appl
5	572	87.2	128	3	US-10-287-994-60	Sequence 60, Appl
6	572	87.2	128	3	US-10-410-997-60	Sequence 60, Appl
7	572	87.2	128	3	US-10-410-962-60	Sequence 60, Appl
8	572	87.2	128	3	US-10-410-897A-60	Sequence 60, Appl
9	572	87.2	128	3	US-10-410-945A-60	Sequence 60, Appl
10	572	87.2	128	3	US-10-410-930A-60	Sequence 60, Appl
11	572	87.2	128	3	US-10-411-012-60	Sequence 60, Appl
12	572	87.2	128	3	US-10-410-913A-60	Sequence 60, Appl
13	572	87.2	128	3	US-11-404-266-60	Sequence 60, Appl
14	572	87.2	128	3	US-10-411-049-60	Sequence 60, Appl
15	566	86.3	128	2	US-09-724-138-46	Sequence 46, Appl
16	566	86.3	128	2	US-09-630-198-46	Sequence 46, Appl
17	559	85.2	129	1	US-08-449-287-2	Sequence 2, Appli
18	559	85.2	235	2	US-09-423-439-18	Sequence 18, Appl
19	559	85.2	235	2	US-09-423-439-58	Sequence 58, Appl
20	559	85.2	235	2	US-09-011-769A-23	Sequence 23, Appl
21	558	85.1	235	2	US-09-238-741-2	Sequence 2, Appli
22	548	83.5	128	2	US-08-619-491-2	Sequence 2, Appli
23	548	83.5	128	5	PCT-US95-07302-2	Sequence 2, Appli
24	543	82.8	128	1	US-07-946-421-26	Sequence 26, Appl
25	543	82.8	235	1	US-08-303-569B-5	Sequence 5, Appli
26	543	82.8	235	1	US-08-116-247-5	Sequence 5, Appli
27	543	82.8	235	2	US-09-795-515-5	Sequence 5, Appli
28	543	82.8	235	2	US-09-348-224-5	Sequence 5, Appli
29	543	82.8	235	3	US-10-704-352-5	Sequence 5, Appli
30	543	82.8	235	3	US-10-704-071-5	Sequence 5, Appli
31	543	82.8	235	3	US-10-703-963-5	Sequence 5, Appli
32	543	82.8	235	3	US-10-703-344-5	Sequence 5, Appli
33	537	81.9	130	3	US-11-143-737-50	Sequence 50, Appl
34	532	81.1	129	1	US-08-116-778E-2	Sequence 2, Appli
35	532	81.1	129	1	US-08-438-562-2	Sequence 2, Appli
36	532	81.1	129	1	US-08-483-528B-92	Sequence 92, Appl
37	531	80.9	128	1	US-08-656-586-2	Sequence 2, Appli
38	524	79.9	128	1	US-07-634-278-31	Sequence 31, Appl
39	524	79.9	128	1	US-08-477-728-31	Sequence 31, Appl
40	524	79.9	128	1	US-08-474-040-31	Sequence 31, Appl
41	524	79.9	128	1	US-08-487-200-31	Sequence 31, Appl
42	524	79.9	128	2	US-08-484-537-31	Sequence 31, Appl
43	524	79.9	128	2	US-09-453-718B-87	Sequence 87, Appl
44	524	79.9	128	3	US-09-718-998-31	Sequence 31, Appl
45	524	79.9	128	3	US-10-160-232-87	Sequence 87, Appl

ALIGNMENTS

RESULT 1

US-08-476-275-4

; Sequence 4, Application US/08476275

; Patent No. 5776456

; GENERAL INFORMATION:

```

; APPLICANT: Anderson, Darrell R.
; APPLICANT: Hanna, Nabil
; APPLICANT: Leonard, John E.
; APPLICANT: Newman, Roland A.
; APPLICANT: Reff, Mitchell E.
; APPLICANT: Rastetter, William H.
; TITLE OF INVENTION: Therapeutic Application of Chimeric and
; TITLE OF INVENTION: Radiolabeled Antibodies to Human B Lymphocyte Restricted
; TITLE OF INVENTION: Differentiation Antigen for the Treatment of B-Cell
; TITLE OF INVENTION: Lymphoma
; NUMBER OF SEQUENCES: 11
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS
; STREET: 699 Prince St.
; CITY: Alexandria
; STATE: VA
; COUNTRY: USA
; ZIP: 22314
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/476,275
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/149,099
; FILING DATE: 03-NOV-1993
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/978,891
; FILING DATE: 13-NOV-1992
; ATTORNEY/AGENT INFORMATION:
; NAME: Teskin, Robin L.
; REGISTRATION NUMBER: 35,030
; REFERENCE/DOCKET NUMBER: 012712-155
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 703-836-6620
; TELEFAX: 703-836-2021
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 128 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-476-275-4

```

```
Query Match      87.2%;  Score 572;  DB 1;  Length 128;
Best Local Similarity  89.8%;  Pred. No. 6e-50;
Matches 114;  Conservative    2;  Mismatches 11;  Indels    0;  Gaps    0;
```

Qy	1	MDFQVQIFSFLLISASVIMSRGQIVLSQSPAILFASPGETVTMTCRASSSVIYMCWNQQK	60
		:	
Db	1	MDFQVQIISFLLISASVIMSRGQIVLSQSPAILSASPGEKVTMTCRASSSVSYIHWFQQK	60
Qy	61	PGSSPKPWIYGTSTLASGVPTRFSGSGSGTSYSLTISRVEAEDAATYYCQOWSSNPFTFG	120

US-08-475-815B-7

; Patent No. 6399061

; GENERAL INFORMATION:

APPLICANT: HANNA, NABIL

; APPLICANT: LEONARD, JOHN E.

; APPLICANT: NEWMAN, ROLAND A.

APPLICANT: REFF, MITCHELL E.

APPLICANT: RASTETTER, WILLIAM H.

; TITLE OF INVENTION: THERAPEUTIC APPLICATION OF CHIMERIC AND

; TITLE OF INVENTION: RADIOLABELED ANTIBODIES TO HUMAN B LYMPHOCYTE RESTRICTED

; TITLE OF INVENTION: DIFFERENTIATION ANTIGEN FOR THE TREATMENT OF B CELL

; TITLE OF INVENTION: LYMPHOMA

```
; NUMBER OF SEQUENCES: 11
```

CORRESPONDENCE ADDRESS:

; ADDRESSEE: PILLSBURY WINTHROP

STREET: 1100 New York Avenue, N.W., Ninth FL.

; CITY: Washington

```
; STATE: DC
```

; COUNTRY: USA

; ZIP: 20005

; COMPUTER READABLE FORM:

```
; MEDIUM TYPE: Floppy disk
```

```
;      COMPUTER:  IBM PC compatible
```

```
; OPERATING SYSTEM: PC-DOS/MS-DOS
```

```
; SOFTWARE: PatentIn Release #1.0, Version #1.30
```

```
; CURRENT APPLICATION DATA:
```

APPLICATION NUMBER: US/08/475,815B

; FILING DATE: 07-JUN-1995

CLASSIFICATION:

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: US 08/149,099

; FILING DATE: 03-NOV-1993

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: US 07/978,891

; FILING DATE: 13-NOV-1992

; ATTORNEY/AGENT INFORMATION:

```
;      NAME:  Teskin, Robin L.
```

; REGISTRATION NUMBER: 35,030

REFERENCE/DOCKET NUMBER: 23522-0157

; TELECOMMUNICATION INFORMATION:

TELEPHONE: 202-861-3000

; TELEFAX: 202-822-0944

; INFORMATION FOR SEQ ID NO: 7:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 128 amino acids

```
;      TYPE:  amino acid
```

Query Match 87.2%; Score 572; DB 2; Length 128;
Best Local Similarity 89.8%; Pred. No. 6e-50;
Matches 114; Conservative 2; Mismatches 11; Indels 0; Gaps 0;

Qy	1	MDFQVQIFSFLLISASVIMSRGQIVLSQSPAILFASPGETVTMTCRASSSVIYMCWNQQK	60
Db	1	MDFQVQIISFLLISASVIMSRGQIVLSQSPAILSASPGEKVTMTCRASSSVSYIHWFQQK	60
Qy	61	PGSSPKPWIYGTSTLASGVPTRFSGSGSGTSYSLTISRVEAEDAATYYCQQWSSNPFTFG	120
		:	
Db	61	PGSSPKPWIYATSNLASGVPVRFSGSGSGTSYSLTISRVEAEDAATYYCQQWTSNPPTFG	120
Qy	121	SGTKLEI	127
Db	121	GGTKLEI	127

US-08-475-813-4

http://es/ScoreAccessWeb/GetItem.action?AppId=106870...0_104727_us-10-687-035-33.rai&ItemType=4&startByte=0 (5 of 18)3/28/2008 11:27:48 AM

Query Match 87.2%; Score 572; DB 2; Length 128;
Best Local Similarity 89.8%; Pred. No. 6e-50;
Matches 114; Conservative 2; Mismatches 11; Indels 0; Gaps 0;

Qy	1	MDFQVQI F S F L L I S A S V I M S R G Q I V L S Q S P A I L F A S P G E T V T M T C R A S S S V I Y M C W N Q Q K	60
		:	
Db	1	MDFQVQII S F L L I S A S V I M S R G Q I V L S Q S P A I L S A S P G E K V T M T C R A S S S V S Y I H W F Q Q K	60
Qy	61	P G S S P K P W I Y G T S T L A S G V P T R F S G S G S G T S Y S L T I S R V E A E D A A T Y Y C Q Q W S S N P P T F G	120
		:	
Db	61	P G S S P K P W I Y A T S N L A S G V P V R F S G S G S G T S Y S L T I S R V E A E D A A T Y Y C Q Q W T S N P P T F G	120
Qy	121	S G T K L E I	127
Db	121	G G T K L E I	127

```
; Sequence 60, Application US/10411037
; Patent No. 7125843
; GENERAL INFORMATION:
; APPLICANT: Neose Technologies, Inc.
; APPLICANT: DeFrees, Shawn
; APPLICANT: Zopf, David
; APPLICANT: Bayer, Robert
; APPLICANT: Hakes, David
; APPLICANT: Chen, Xi
; APPLICANT: Bowe, Caryn
; TITLE OF INVENTION: ALPHA GALACTOSIDASE A: REMODELING AND GLYCOCONJUGATION OF ALPHA
; TITLE OF INVENTION: GALACTOSIDASE A
; FILE REFERENCE: 040853-01-5082
; CURRENT APPLICATION NUMBER: US/10/411,037
; CURRENT FILING DATE: 2003-04-09
; PRIOR APPLICATION NUMBER: US 60/328,523
; PRIOR FILING DATE: 2001-10-10
; PRIOR APPLICATION NUMBER: US 60/344,692
; PRIOR FILING DATE: 2001-10-19
; PRIOR APPLICATION NUMBER: US 60/387,292
; PRIOR FILING DATE: 2002-06-07
; PRIOR APPLICATION NUMBER: US 60/391,777
```

Query Match 87.2%; Score 572; DB 3; Length 128;
Best Local Similarity 89.8%; Pred. No. 6e-50;
Matches 114; Conservative 2; Mismatches 11; Indels 0; Gaps 0;

Qy	1	MDFQVQIIFSLLLISASVIMSRGQIVLSQSPAILFASPGETVTMTCRASSSVIYMCWNQQK	60
Db	1	MDFQVQIISFLLISASVIMSRGQIVLSQSPAILSASPGEKVTMTCRASSSVSYIHWFQQK	60
Qy	61	PGSSPKPWIIYGTSTLASGVPTRFSGSGSGTSYSLTISRVEAEDAATYYCQQWSSNPFTFG	120
Db	61	PGSSPKPWIIYATSNLASGVPVRFSGSGSGTSYSLTISRVEAEDAATYYCQQWTSNPFTFG	120
Qy	121	SGTKLEI	127
Db	121	GGTKLEI	127

```
US-10-287-994-60
; Sequence 60, Application US/10287994
; Patent No. 7138371
; GENERAL INFORMATION:
; APPLICANT: Neose Technologies, Inc.
; APPLICANT: DeFrees, Shawn
; APPLICANT: Zopf, David
; APPLICANT: Bayer, Robert
; APPLICANT: Bowe, Caryn
; APPLICANT: Hakes, David
; APPLICANT: Chen, Xi
; TITLE OF INVENTION: REMODELING AND GLYCOCONJUGATION OF PEPTIDES
; FILE REFERENCE: 040853-01-5052-00
; CURRENT APPLICATION NUMBER: US/10/287,994
; CURRENT FILING DATE: 2002-11-05
; PRIOR APPLICATION NUMBER: US 60/328,523
; PRIOR FILING DATE: 2001-10-10
; PRIOR APPLICATION NUMBER: US 60/344,692
; PRIOR FILING DATE: 2001-10-19
; PRIOR APPLICATION NUMBER: US 60/387,292
; PRIOR FILING DATE: 2002-06-07
; PRIOR APPLICATION NUMBER: US 60/391,777
; PRIOR FILING DATE: 2002-06-25
; PRIOR APPLICATION NUMBER: US 60/396,594
```

```
; PRIOR FILING DATE: 2002-07-17
; PRIOR APPLICATION NUMBER: US 60/404,249
; PRIOR FILING DATE: 2002-08-16
; PRIOR APPLICATION NUMBER: US 60/407,527
; PRIOR FILING DATE: 2002-08-28
; NUMBER OF SEQ ID NOS: 62
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 60
;   LENGTH: 128
;   TYPE: PRT
;   ORGANISM: Mus musculus
US-10-287-994-60
```

Query Match 87.2%; Score 572; DB 3; Length 128;
Best Local Similarity 89.8%; Pred. No. 6e-50;
Matches 114; Conservative 2; Mismatches 11; Indels 0; Gaps 0;

Qy	1	MDFQVQIFSFLLISASVIMSRGQIVLSQSPAILFASPGETVTMTCRASSSVIYMCWNQQK	60
Db	1	MDFQVQIISFLLISASVIMSRGQIVLSQSPAILSASPGEKVTMTCRASSSVSYIHWFQQK	60
Qy	61	PGSSPKPWIYGTSTLASGVPTRFSGSGSGTSYSLTISRVEAEDAATYYCQQWSSNPFTFG	120
Db	61	PGSSPKPWIYATSNLASGVPVRFSGSGSGTSYSLTISRVEAEDAATYYCQQWTSNPFTFG	120
Qy	121	SGTKLEI	127
Db	121	GGTKLEI	127

RESULT 6

```
US-10-410-997-60
; Sequence 60, Application US/10410997
; Patent No. 7157277
; GENERAL INFORMATION:
; APPLICANT: Neose Technologies, Inc.
; APPLICANT: DeFrees, Shawn
; APPLICANT: Zopf, David
; APPLICANT: Bayer, Robert
; APPLICANT: Hakes, David
; APPLICANT: Chen, Xi
; APPLICANT: Bowe, Caryn
; TITLE OF INVENTION: FOLLICLE STIMULATING HORMONE: REMODELING AND GLYCOCONJUGATION OF
; TITLE OF INVENTION: FSH
; FILE REFERENCE: 040853-01-5059
; CURRENT APPLICATION NUMBER: US/10/410,997
; CURRENT FILING DATE: 2003-04-09
; PRIOR APPLICATION NUMBER: US 60/328,523
; PRIOR FILING DATE: 2001-10-10
; PRIOR APPLICATION NUMBER: US 60/344,692
; PRIOR FILING DATE: 2001-10-19
; PRIOR APPLICATION NUMBER: US 60/387,292
; PRIOR FILING DATE: 2002-06-07
; PRIOR APPLICATION NUMBER: US 60/391,777
; PRIOR FILING DATE: 2002-06-25
; PRIOR APPLICATION NUMBER: US 60/396,594
; PRIOR FILING DATE: 2002-07-17
```


Query Match 87.2%; Score 572; DB 3; Length 128;
Best Local Similarity 89.8%; Pred. No. 6e-50;
Matches 114; Conservative 2; Mismatches 11; Indels 0; Gaps 0;

```

RESULT 7
US-10-410-962-60
; Sequence 60, Application US/10410962
; Patent No. 7173003
; GENERAL INFORMATION:
; APPLICANT: Neose Technologies, Inc.
; APPLICANT: DeFrees, Shawn
; APPLICANT: Zopf, David
; APPLICANT: Bayer, Robert
; APPLICANT: Hakes, David
; APPLICANT: Chen, Xi
; APPLICANT: Bowe, Caryn
; TITLE OF INVENTION: GRANULOCYTE COLONY STIMULATING FACTOR: REMODELING AND
; TITLE OF INVENTION: GLYCOCONJUGATION OF G-CSF
; FILE REFERENCE: 040853-01-5054
; CURRENT APPLICATION NUMBER: US/10/410,962
; CURRENT FILING DATE: 2003-04-09
; PRIOR APPLICATION NUMBER: US 60/328,523
; PRIOR FILING DATE: 2001-10-10
; PRIOR APPLICATION NUMBER: US 60/344,692
; PRIOR FILING DATE: 2001-10-19
; PRIOR APPLICATION NUMBER: US 60/387,292
; PRIOR FILING DATE: 2002-06-07
; PRIOR APPLICATION NUMBER: US 60/391,777
; PRIOR FILING DATE: 2002-06-25
; PRIOR APPLICATION NUMBER: US 60/396,594
; PRIOR FILING DATE: 2002-07-17
; PRIOR APPLICATION NUMBER: US 60/404,249

```

Query Match 87.2%; Score 572; DB 3; Length 128;
Best Local Similarity 89.8%; Pred. No. 6e-50;
Matches 114; Conservative 2; Mismatches 11; Indels 0; Gaps 0;

Qy	1	MDFQVQIFSFLLISASVIMSRGQIVLSQSPAILFASPGETVTMTCRASSSVIYMCWNQQK	60
Db	1	MDFQVQIISFLLISASVIMSRGQIVLSQSPAILSASPGEKVTMTCRASSSVSYIHWFQQK	60
Qy	61	PGSSPKPWIYGTSTLASGVPTRFSGSGSGTSYSLTISRVEAEDAATYYCQQWSSNPFTFG	120
Db	61	PGSSPKPWIYATSNLASGVPVRFSGSGSGTSYSLTISRVEAEDAATYYCQQWTSNPPTFG	120
Qy	121	SGTKLEI	127
Db	121	GGTKLEI	127

US-10-410-897A-60

http://es/ScoreAccessWeb/GetItem.action?AppId=106870...0_104727_us-10-687-035-33.rai&ItemType=4&startByte=0 (10 of 18)3/28/2008 11:27:48 AM

Query Match 87.2%; Score 572; DB 3; Length 128;
Best Local Similarity 89.8%; Pred. No. 6e-50;
Matches 114; Conservative 2; Mismatches 11; Indels 0; Gaps 0;

Qy	1	MDFQVQIFSFLLISASVIMSRGQIVLSQSPAILFASPGETVTMTCRASSSVIYMCWNQQK	60
Db	1	MDFQVQIISFLLISASVIMSRGQIVLSQSPAILSASPGEKVTMTCRASSSVSYIHWFQQK	60
Qy	61	PGSSPKPWIYGTSTLASGVPTRFSGSGSGTSYSLTISRVEAEDAATYYCQQWSSNPFTFG	120
Db	61	PGSSPKPWIYATSNLASGVPVRFSGSGSGTSYSLTISRVEAEDAATYYCQQWTSNPFTFG	120
Qy	121	SGTKLEI	127
Db	121	GGTKLEI	127

US-10-410-945A-60

http://es/ScoreAccessWeb/GetItem.action?AppId=106870...0_104727_us-10-687-035-33.rai&ItemType=4&startByte=0 (11 of 18)3/28/2008 11:27:48 AM

Query Match 87.2%; Score 572; DB 3; Length 128;
Best Local Similarity 89.8%; Pred. No. 6e-50;
Matches 114; Conservative 2; Mismatches 11; Indels 0; Gaps 0;

Qy	1	MDFQVQIFSFLLISASVIMSRGQIVLSQSPAILFASPGETVTMTCRASSSVIYMCWNQQK	60
		:	
Db	1	MDFQVQIIISFLLISASVIMSRGQIVLSQSPAILSASPGEKVMTTCRASSSVSYIHWFQQK	60
Qy	61	PGSSPKPWIIYGTSTLASGVPTRFSGSGSGTSYSLTISRVEAEDAATYYCQQWSSNPFTFG	120
		:	
Db	61	PGSSPKPWIYATSNLASGPVVRFSGSGSGTSYSLTISRVEAEDAATYYCQQWTSNPPTFG	120
Qy	121	SGTKLEI	127
Db	121	GGTKLEI	127

US-10-410-930A-60

```
; Sequence 60, Application US/10410930A
; Patent No. 7226903
```

; GENERAL INFORMATION:

; APPLICANT: Neose Technologies, Inc.
; APPLICANT: DeFrees, Shawn
; APPLICANT: Zopf, David
; APPLICANT: Bayer, Robert
; APPLICANT: Hakes, David
; APPLICANT: Chen, Xi
; APPLICANT: Bowe, Caryn

```
; TITLE OF INVENTION: INTERFERON BETA: REMODELING AND GLYCOCONJUGATION OF INTERFERON BETA
; FILE REFERENCE: 40853-01-5056-US
; CURRENT APPLICATION NUMBER: US/10/410,930A
; CURRENT FILING DATE: 2003-04-09
; PRIOR APPLICATION NUMBER: US 60/328,523
; PRIOR FILING DATE: 2001-10-10
; PRIOR APPLICATION NUMBER: US 60/344,692
```

```

Query Match      87.2%;   Score 572;   DB 3;   Length 128;
Best Local Similarity 89.8%;   Pred. No. 6e-50;
Matches 114;   Conservative 2;   Mismatches 11;   Indels 0;   Gaps 0;

Qy      1 MDFQVQIFSFL LISASVIMSRGQIVLSQSPAILFASPGETVTMTCRASSSVIYMCWNQQK 60
      ||||| ||||||||||||||||||||| ||||| ||||||||||| : | |||
Db      1 MDFQVQIISFLLISASVIMSRGQIVLSQSPAILSASPGEKVTMTCRASSSVSYIHWFQQK 60

Qy      61 PGSSPKPWIYGTSTLASGVPTRFSGSGSGTSYSLTISRVEAEDAATYYCQQWSSNPFTFG 120
      ||||| || ||||| ||||||||||||||||||||||||||||| : ||| |||
Db      61 PGSSPKPWIYATSNLASGVPVRFSGSGSGTSYSLTISRVEAEDAATYYCQQWTSNPPTFG 120

Qy      121 SGTKLEI 127
      |||||
Db      121 GGTKLEI 127

```

http://es/ScoreAccessWeb/GetItem.action?AppId=106870...0_104727_us-10-687-035-33.rai&ItemType=4&startByte=0 (13 of 18)3/28/2008 11:27:48 AM

```

Query Match      87.2%;   Score 572;   DB 3;   Length 128;
Best Local Similarity  89.8%;   Pred. No. 6e-50;
Matches 114;   Conservative      2;   Mismatches 11;   Indels      0;   Gaps      0;

Qy      1 MDFQVQIFSFLLISASVIMSRGQIVLSQSPAILFASPGETVTMTCRASSSVIYMCWNQQK 60
      ||||| ||||||||||||||||||||||| |||| | ||||||||| | : | |||
Db      1 MDFQVQIISFLLISASVIMSRGQIVLSQSPAILSASPGEKVTMTCRASSSVSYIHWFQQK 60

Qy      61 PGSSPKPWIYGTSTLASGVPTRFSGSGSGTSYSLTISRVEAEDAATYYCQQWSSNPFTFG 120
      ||||| || || ||||| ||||||||||||||||||||||||||||| : ||| |||
Db      61 PGSSPKPWIYATSNLASGVPVRFSGSGSGTSYSLTISRVEAEDAATYYCQQWTSNPPTFG 120

Qy      121 SGTKLEI 127
      |||||
Db      121 GGTKLEI 127

```

US-10-410-913A-60

http://es/ScoreAccessWeb/GetItem.action?AppId=106870...0_104727_us-10-687-035-33.rai&ItemType=4&startByte=0 (14 of 18)3/28/2008 11:27:48 AM

RESULT 13
US-11-404-266-60
; Sequence 60, Application US/11404266
; Patent No. 7276475
; GENERAL INFORMATION:
; APPLICANT: Neose Technologies, Inc.
; APPLICANT: DeFrees, Shawn
; APPLICANT: Zopf, David
; APPLICANT: Bowe, Caryn

RESULT 14
US-10-411-049-60
; Sequence 60, Application US/10411049
; Patent No. 7297511
; GENERAL INFORMATION:
; APPLICANT: Neose Technologies, Inc.
; APPLICANT: DeFrees, Shawn

Query Match 86.3%; Score 566; DB 2; Length 128;
Best Local Similarity 88.2%; Pred. No. 2.4e-49;
Matches 112; Conservative 4; Mismatches 11; Indels 0; Gaps 0;

Qy	1	MDFQVQIFSFLLISASVIMSRGQIVLSQSPAILFASPGETVTMTCRASSSVIYMCWNQQK	60
		:	
Db	1	MDFQVQIFSFLLISASVIIARGQIVLSQSPAILSASPGEKVTMTCRASSSVSYMHWYQQK	60
Qy	61	PGSSPKPWIYGTSTLASGVPTRFSGSGGTSYSLTISRVEAEDAATYYCQQWSSNPFTFG	120
Db	61	PGSSPKPWIYAPSNLASGVPARFSGSGGTSYSLTISRVEAEDAATYYCQQWSFNPPFTFG	120
Qy	121	SGTKLEI	127
		: :	
Db	121	AGTKLEL	127

Search completed: March 10, 2008, 14:05:32
Job time : 37.8083 secs